```
SEQUENCE LISTING
(1) GENERAL INFORMATION:
        (i) APPLICANT: Kulesz-Martin, Molly F.
        (ii) TITLE OF INVENTION: p53as PROTEIN AND ANTIBODY THEREFOR
        (iii) NUMBER OF SEQUENCES: 5
        (iv) CORRESPONDENCE ADDRESS:
                 (A) ADDRESSEE: Dunn & Simpson, P.C.
                      STREET: P.O. Box 96
                 (C) CITY: Newfane
                 (D) STATE: New York
                     COUNTRY: U.S.A.
                 (F) ZIP: 14108
        (v) COMPUTER READABLE FORM:
                 (A) MEDIUM TYPE: Diskette - 3/50 inch, 1.44 Mb storage
                 (B) COMPUTER: Victor 300 SX//25 (IBM PC Compatible)
                 (C) OPERATING SYSTEM: MS-DOS/Version 5.0
                 (D) SOFTWARE: Wordstar Prof∉ssional Release 4
        (vi) CURRENT APPLICATION DATA:
                (A) APPLICATION NUMBER: 08/100,496
                (B) FILING DATE: 2-Aug-1993
                (C) CLASSIFICATION: 530
        (vii) PRIOR APPLICATION DATA:
                (A) APPLICATION NUMBER:
                (B) FILING DATE:
        (viii) ATTORNEY/AGENT INFORMATION:
                (A) NAME: Dunn, Michael L.
                (B) REGISTRATION NUMB/ER:
                                          25,330
                (C) REFERENCE/DOCKET/NUMBER: RPP:135 US
        (ix) TELECOMMUNICATION INFORMATION:
                (A) TELEPHONE (716)/433-1661
                (B) TELEFAX: (716)/433-1665
(2) INFORMATION FOR SEQ ID NO
        (i) SEQUENCE CHARACTERISTICS:
                (A) LENGTH: 17
                     TYPE: amin/o acids
                (B)
                (C) STRANDEDNESS: n/a
                (D) TOPOLOGY: /n/a
        (ii) MOLECULE TYPE: peptide
        (iii) HYPOTHETICAL: no
        (iv) ANTI-SENSE: no
        (v) FRAGMENT TYPE: /h/a
        (vi) ORIGINAL SOURCE:
                (A) ORGANISM: mouse
                (B) STRAIN: n/a
                (C) INDIVIDUAL ISOLATE:
                (D) DEVELOPMENTAL STAGE: n/a
                (E) HAPLOTYPE: n/a
                (F) TLSSUE TYPE: n/a
                (G) CELL TYPE: n/a
                (H) ÇELL LINE: n/a
                (I) ∕ORGANELLE: n/a
        (vii) IMMEDIATE SOURCE: sequenced from cDNA clone from mouse
                   epidermal cell RNA, Genbank Accession #M13874
                (A) LIBRARY: plasmid p6.3
                (B) CLONE:
        (viii) POSITION IN GENOME:
```

(ix) FEATURE: n/a

(C) UNITS:

/(A) CHROMOSOME/SEGMENT: 1 (B) MAP POSITION: p53 gene

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(A) NAME/KEY:
                 (B) LOCATION:
                 (C) IDENTIFICATION METHOD:
                 (D) OTHER INFORMATION:
         (x) PUBLICATION INFORMATION:
                 (A) AUTHORS: Kulesz-Martin et al.
                 (B) TITLE: Endogenous p53 Protein Generated From Wild
                     Type Alternatively Spliced P5/3 RNA in Mouse
                · (C) JOURNAL: Mol. Cell. Biol.
                 (D) VOLUME: 14
                 (E) ISSUE: 3
                 (F) PAGES: 1698-1708
                 (G) DATE: March, 1994
                 (A) AUTHORS: Han, K.A. and Kul/esz-Martin, M.F.
                 (B) TITLE: Alternatively Spliced p53 RNA in Transformed
                     and Normal Cells of Different Tissue Types
                 (C) JOURNAL: Nucleic Acids Res.
                 (D) VOLUME: 20
                 (E) ISSUE: 8
                 (F) PAGES: 1979-1981
                 (G) DATE: 1992
                 (A) AUTHORS:
                               Arai, N. et al.
                 (B) TIŢLE: Immunological/ly Distinct p53 Molecules Generated
                     by Alternative Spliging
                 (C) JOURNAL: Mol. and Gell. Biol.
                 (D) VOLUME: 6
                 (E) ISSUE:
                 (F) PAGES: 3232-3239
                 (G) DATE: 1986
                 (H) DOCUMENT NUMBER
                 (I) FILING DATE:
                 (J) PUBLICATION DATE:
                 (K) RELEVANT RESIDUES IN SEQ ID NO:
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1
Leu Gln Pro Arg Ala Phe Gln Ala Leu Ile Lys Glu Glu Ser Pro Asn
                 5
                                      10
Cys
(3) INFORMATION FOR SEQ ID / NO:
       (i) SEQUENCE CHARACT/ERISTICS:
                (A) LENGTH:/33
                    TYPE: /Nucleic Acids
                (C) STRANDEDNESS: Unknown
                (D) TOPOLØGY: Unknown
        (ii) MOLECULE TYPE: Oligonucleotide
        (iii) HYPOTHETIÇAL:
        (iv) ANTI-SENSE
        (V) FRAGMENT TYPE:
        (vi) ORIGINAL SOURCE:
                (A) OAGANISM: Human
                    $TRAIN:
                (B)
                (C)
                    /INDIVIDUAL ISOLATE:
                (D)/DEVELOPMENTAL STAGE:
                (E) HAPLOTYPE:
                (月) TISSUE TYPE:
                (Ġ) CELL TYPE:
                (H) CELL LINE:
                (I) ORGANELLE:
        (Vii) LMMEDIATE SOURCE: Genbank Accession #X54156, Locus HUM P53G
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(A) LIBRARY:
                 (B)
                      CLONE
        (viii) POSITION IN GENOME:
                 (A) CHROMOSOME/SEGMENT: 17
                 (B) MAP POSITION: p53 gene 17593-17613
                 (C) UNITS:
        (ix) FEATURE:
                 (A) NAME/KEY:
                 (B) LOCATION:
                 (C) IDENTIFICATION METHOD:
                 (D) OTHER INFORMATION:
        (x) PUBLICATION INFORMATION:
                 (A) AUTHORS:
                 (B) TITLE:
                 (C) JOURNAL:
                 (D) VOLUME:
                 (E) ISSUE:
                 (F) PAGES:
                 (G) DATE:
                 (H) DOCUMENT NUMBER:
                 (I) FILING DATE:
                 (J) PUBLICATION DATE:
                 (K) RELEVANT RESIDUES IN SEQ
                                              NO: 2
        (xi) SEQUENCE DESCRIPTION: SEQ
ATCGAAGCTT GAGATGTTCC GAGAGAGCTG AAT
                                              33
(4) INFORMATION FOR SEQ ID
                              NO: 3:
        (i) SEQUENCE CHARACTERISTICS:
                 (A) LENGTH: 31
                     TYPE: Nucleic a/cids
                 (C) STRANDEDNESS: whknown
                 (D) TOPOLOGY: unknown
        (ii) MOLECULE TYPE: oligorucleotide
        (iii) HYPOTHETICAL:
        (iv) ANTI-SENSE:
        (v) FRAGMENT TYPE:
        (vi) ORIGINAL SOURCE:
                 (A) ORGANISM: /Human
                 (B) STRAIN:
                 (C) INDIVIDUAL ISOLATE:
                 (D) DEVELOPMENTAL STAGE:
                 (E) HAPLOTYPE:
                 (F) TISSUE TYPE:
                 (G) CELL/TYPE:
                 (H) CELL LINE:
                 (I) ORGANELLE:
        (vii) IMMEDIATÉ SOURCE: Genbank Accession #54156, Locus HUMP53G
                 (A) LIBRARY:
                 (B)
                     /CLONE
        (viii) POSITION IN GENOME:
                 (A)/ CHROMOSOME/SEGMENT: 17
                 (B) MAP POSITION: p53 gene 18774-18794
                 (¢) UNITS:
        (ix) FEATURE:
                 (A) NAME/KEY:
                 (B) LOCATION:
                 (C) IDENTIFICATION METHOD:
                 (D) OTHER INFORMATION:
             UBLICATION INFORMATION:
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(A) AUTHORS:
                 (B) TITLE:
                 (C) JOURNAL:
                 (D) VOLUME:
                 (E) ISSUE:
                 (F) PAGES:
                 (G) DATE:
                 (H) DOCUMENT NUMBER:
                 (I) FILING DATE:
                 (J) PUBLICATION DATE:
                 (K) RELEVANT RESIDUES IN SEQ
                                                 ΙD
        (xi) SEQUENCE DESCRIPTION: SEQ
                                          ID
                                              NO: 3
ATCGTCTAGA GCTTCTGACG CACACCTATT G
                                                 31
(5) INFORMATION FOR SEQ ID NO: 4:
        (i) SEQUENCE CHARACTERISTICS:
                 (A) LENGTH: 20
                 (B)
                      TYPE: Amino Acids
                 (C) STRANDEDNESS: unknown
                 (D) TOPOLOGY: unknown
        (ii) MOLECULE TYPE: Peptide
        (iii) HYPOTHETICAL: deduced from intron 10 sequences p53 gene
        (iv) ANTI-SENSE:
        (V) FRAGMENT TYPE:
        (vi) ORIGINAL SOURCE:
                 (A) ORGANISM: Human
                 (B) STRAIN:
                 (C) INDIVIDUAL ISOLATE:
                 (D) DEVELOPMENTAL STAGE
                 (E) HAPLOTYPE:
                 (F) TISSUE TYPE:
                 (G) CELL TYPE:
                 (H) CELL LINE:
                 (I) ORGANELLE:
        (vii) IMMEDIATE SOURCE:
                 (A) LIBRARY: deducéd translation from nucleotides in
                     Genbank nucleic acid database accession #54156,
                     Locus HSP53G
                    CLONE:
                 (B)
        (viii) POSITION IN GENOME:
                (A) CHROMOSOME/SEGMENT:
                                           17
                 (B) MAP POSITION: p53 gene, at 18530 to 18589
                 (C) UNITS:
        (ix) FEATURE: n/a
                (A) NAME/KEY:
                (B) LOCATION:
                (C) IDENTIFICATION METHOD:
                (D) OTHER/INFORMATION:
        (x) PUBLICATION INFORMATION:
                (A) AUTHORS:
                (B) TITLE:
                (C) JOURNAL:
                (D) VØLUME:
                (E)
                    ISSUE:
                (F)
                    /PAGES:
                (G)/
                    DATE:
                (H) DOCUMENT NUMBER:
                    FILING DATE:
                    PUBLICATION DATE:
                                       53
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(K) RELEVANT RESIDUES IN SEQ
         (xi) SEQUENCE DESCRIPTION: SEQ
                                          ID NO: 4
Arg Glu Lys Gly His Arg Pro Ser His Ser Cys Asp Val Il¢ Ser Pro
                                       10
Pro Cys Phe Cys
              20
(6) INFORMATION FOR SEQ
                          ID
                              NO: 5:
         (i) SEQUENCE CHARACTERISTICS:
                 (A) LENGTH: 16
                 (B)
                      TYPE: Amino Acids
                 (C) STRANDEDNESS: unknown
                 (D) TOPOLOGY: unknown
         (ii) MOLECULE TYPE: peptide
         (iii) HYPOTHETICAL:
         (iv) ANTI-SENSE:
         (V) FRAGMENT TYPE:
         (vi) ORIGINAL SOURCE: mouse glucocoxticoid receptor sequence
                 (A) ORGANISM:
                 (B) STRAIN:
                 (C) INDIVIDUAL ISOLATE:
                 (D) DEVELOPMENTAL STAGE:
                 (E) HAPLOTYPE:
                 (F) TISSUE TYPE:
                 (G) CELL TYPE:
                 (H) CELL LINE:
                 (I) ORGANELLE:
         (vii) IMMEDIATE SOURCE:
                 (A) LIBRARY:
                 (B)
                      CLONE
         (viii) POSITION IN GENOME:
                 (A) CHROMOSOME/SEGMENT:
                 (B) MAP POSITION:
                 (C) UNITS:
        (ix) FEATURE:
                 (A) NAME/KEY:
                 (B) LOCATION:
                 (C) IDENTIFICATION METHOD:
                 (D) OTHER INFORMATION:
        (x) PUBLICATION INFORMATION:
                 (A) AUTHORS
                 (B) TITLE:
                 (C) JOURNAL:
                 (D) VOLUME:
                 (E) ISSUE:
                 (F) PAGES:
                 (G) DATE:
                 (H) DOÇUMENT NUMBER:
                 (I) FILING DATE:
                 (J) PØBLICATION DATE:
                 (K) RELEVANT RESIDUES IN SEQ ID
        (xi) SEQUENÇE DESCRIPTION: SEQ ID NO: 5
Gly Arg Asn Asp Cys/Ile Ile Asp Lys Ile Arg Arg Lys Asn Cys Asp
                                      10
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